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(FILE 'HOME' ENTERED AT 09:55:04 ON 27 FEB 2007)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 09:55:18 ON 27 FEB 2007 SEA (HYDANTOIN RACEMASE)

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FILE AGRICOLA
  4
 9
     FILE BIOENG
22
      FILE BIOSIS
42
     FILE BIOTECHABS
42
      FILE BIOTECHDS
      FILE BIOTECHNO
11
      FILE CABA
 2
      FILE CAPLUS
57
 7
      FILE CEABA-VTB
      FILE CIN
 1
     FILE DGENE
107
15
     FILE EMBASE
     FILE ESBIOBASE
14
     FILE FSTA
 5
      FILE GENBANK
139
      FILE IFIPAT
23
      FILE JICST-EPLUS
 3
      FILE LIFESCI
16
17
      FILE MEDLINE
16
      FILE PASCAL
12
      FILE PCTGEN
      FILE PROMT
 1
      FILE SCISEARCH
25
      FILE TOXCENTER
 7
      FILE USPATFULL
33
      FILE USPAT2
 8
      FILE WPIDS
21
     FILE WPINDEX
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FILE 'CAPLUS, BIOTECHDS, SCISEARCH, IFIPAT, BIOSIS, WPIDS' ENTERED AT 09:58:53 ON 27 FEB 2007

L2 190 S L1

L1

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L3 21 S L2 AND (MUTANT OR VARIANT)

L4 9 DUP REM L3 (12 DUPLICATES REMOVED)

QUE (HYDANTOIN RACEMASE)

L4 ANSWER 1 OF 9 BIOTECHDS COPYRIGHT 2007 THE THOMSON CORP. on STN

DUPLICATE 1

ACCESSION NUMBER: 2006-27154 BIOTECHDS

TITLE: Novel 5-substituted hydantoin racemase

polypeptide useful for racemization of optically active 5-substituted hydantoin, and for producing optically active

N-carbamyl amino acid or amino acid;

involving vector plasmid pBHR001-mediated gene transfer

and expression in Bacillus sp. or Escherichia coli

AUTHOR: NISHI K; YANAGISAWA S; NANBA H; UEDA M; NORO N

PATENT ASSIGNEE: KANEKA CORP

PATENT INFO: WO 2006080409 3 Aug 2006 APPLICATION INFO: WO 2006-JP301253 26 Jan 2006

PRIORITY INFO: JP 2005-22802 31 Jan 2005; JP 2005-22802 31 Jan 2005

DOCUMENT TYPE: Patent LANGUAGE: Japanese

OTHER SOURCE: WPI: 2006-755827 [77]

AB DERWENT ABSTRACT:

NOVELTY - A 5-substituted hydantoin racemase

polypeptide (I) having molecular weight of 139000, Km value of 0.304 mM with respect to L-5-(2-methylthioethyl)hydantoin, range of temperature being 25-65degreesC (optimum temperature being 40degreesC), range of pH being 6-10 (optimum pH being 8-9), temperature stability at 30degreesC or less, and pH stability at 4.5-8.0, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for: (1) DNA encoding (I) having amino acid sequence of SEQ ID Number 1, consisting of base sequence of SEQ ID Number 2 or its homologous base sequence; (2) recombinant plasmid comprising the DNA encoding (I); (3) transformant obtained by transforming host microorganism with the recombinant plasmid; (4) microorganism belonging to Bacillus genus, capable of producing (I); (5) producing (I); (6) racemization method of optically active 5-substituted hydantoin, comprises allowing (I), the transformant or the microorganism having capability of producing (I), to act on an optically active 5-substituted hydantoin compound; (7) producing optically active N-carbamyl amino acid, involves allowing hydantoinase and (I), the transformant or the microorganism having capability of producing (I), to act on 5-substituted hydantoin compound; and (8) producing optically active amino acid, comprises allowing hydantoinase, N-carbamyl amino acid amide hydrolase and (I), the transformant or the microorganism having capability of producing (I), to act on 5-substituted hydantoin compound.

BIOTECHNOLOGY - Preparation (claimed): (I) is produced by culturing the microorganism having capability of producing (I), such that (I) is accumulated in a culture, and extracting (I) from the culture. The microorganism having capability of producing (I) includes the transformant or the microorganism of Bacillus genus. Preferred Polypeptide: (I) is obtained from the microorganism belonging to Bacillus genus such as Bacillus sp. KNK519HR strain (FERM BP-10477). Preferred Plasmid: The recombinant plasmid is plasmid pBHR001. Preferred Transformant: The host microorganism is Escherichia coli, and the transformant is E. coli HB101(pBHR001) (FERM BP-10476). Preferred Microorganism: The microorganism capable of producing (I), is Bacillus sp. KNK519HR strain (FERM BP-10477) or its mutant.

USE - For racemization method of optically active 5-substituted hydantoin, and for manufacturing optically active N-carbamyl amino acid chosen from N-carbamyl-D-leucine, N-carbamyl-D-isoleucine, N-carbamyl-D-valine, N-carbamyl-D-norleucine, N-carbamyl-D-norvaline, N-carbamyl-D-methionine, N-carbamyl-D-cysteine, N-carbamyl-D-phenyl glycine and N-carbamyl-D-4-hydroxyphenyl glycine. For producing optically active amino acid chosen from D-leucine, D-isoleucine, D-valine, D-norleucine, D-norvaline, D-methionine, D-cysteine, D-penicillamine, D-phenylalanine,

D-phenyl glycine and D-4-hydroxyphenyl glycine (all claimed).

ADVANTAGE - (I) enables efficient production of N-carbamyl amino acid or optically active amino acid.

EXAMPLE - Microorganisms such as Bacillus sp. KNK519HR strain (FERM BP-10477) was inoculated to the culture medium comprising glycerol (in weight%) (1), glucose (0.5), potassium dihydrogen phosphate (KH2PO4) (0.454), disodium hydrogen phosphate (Na2HPO4) (0.620), diammonium sulfate ((NH4)2SO4) (0.65), heptahydrate magnesium sulfate (MgSO4.7H2O) (0.05), tetrahydrate manganese chloride (MnCl2.4H2O) (0.002), heptahydrate ferrous sulfate (FeSO4.4H2O) (0.002), dihydrated calcium chloride (CaCl2.2H2O) (0.002), anhydrous sodium citrate (0.032), DL-5-(2-methylthioethyl)hydantoin (0.05), DL-5-methyl hydantoin (0.05) and DL-5-benzyl-hydantoin (0.05), and shaking culture was aerobically was carried out at 30degreesC for 17 hours. To the culture, the sterilized glucose solution was added. After cultivation, the microbial cells were separated by centrifugation, suspended in 50 mM potassium phosphate buffer comprising 1 mM dithiothreitol (DTT), crushed and subjected to centrifugation. After centrifugation, the ammonium sulfate was added to the supernatant liquid (crude-enzyme liquid), and the precipitate was obtained by centrifugation. The precipitate was dissolved in 50 mM potassium phosphate buffer comprising 1 mM DTT, and subjected to dialysis and column chromatography using TSKgel diethylaminoethyl Toyopearl 650 M, to obtain 93% pure hydantoin racemase. The isolated hydantoin racemase was found to exhibit substrate specificity with respect to D-5-substituted hydantoin compound. (51 pages)

ANSWER 2 OF 9 CAPLUS COPYRIGHT 2007 ACS on STN DUPLICATE 2

ACCESSION NUMBER:

2006:1305721 CAPLUS

DOCUMENT NUMBER:

146:179326

TITLE:

Site-directed mutagenesis indicates an important role

of cysteines 76 and 181 in the catalysis of

hydantoin racemase from Sinorhizobium meliloti

AUTHOR (S):

Martinez-Rodriguez, Sergio; Andujar-Sanchez,

Montserrat; Neira, Jose L.; Clemente-Jimenez, Josefa M.; Jara-Perez, Vicente; Rodriguez-Vico, Felipe;

Heras-Vazquez, Francisco J. Las

CORPORATE SOURCE:

Departamento Quimica Fisica, Bioquimica y Quimica Inorganica, Universidad de Almeria, Almeria, 04120,

Spain

SOURCE:

Protein Science (2006), 15(12), 2729-2738

CODEN: PRCIEI; ISSN: 0961-8368

PUBLISHER:

Cold Spring Harbor Laboratory Press

DOCUMENT TYPE:

Journal

LANGUAGE:

English

Hydantoin racemase enzyme plays a crucial role in the reaction cascade known as "hydantoinase process.". In conjunction with a stereoselective hydantoinase and a stereospecific carbamoylase, it allows the total conversion from D,L-5-monosubstituted hydantoins, with a low rate of racemization, to optically pure D- or L-amino acids. Residues Cys-76 and Cys-181 belonging to hydantoin racemase from Sinorhizobium meliloti (SmeHyuA) have been proved to be involved in catalysis. Here, we report biophys. data of SmeHyuA Cys-76 and Cys-181 to alanine mutants, which point toward a two-base mechanism for the racemization of 5-monosubstituted hydantoins. The secondary and the tertiary structure of the mutants were not significantly affected, as shown by CD. Calorimetric and fluorescence expts. have shown that Cys-76 is responsible for recognition and proton retrieval of D-isomers, while Cys-181 is responsible for L-isomer recognition and racemization. This recognition process is further supported by measurements of protein stability followed by chemical denaturation in the presence of the corresponding compound 50

REFERENCE COUNT:

THERE ARE 50 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

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ANSWER 3 OF 9 CAPLUS COPYRIGHT 2007 ACS on STN DUPLICATE 3
                          2006:178846 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                          144:407108
                          Binding studies of hydantoin
TITLE:
                          racemase from Sinorhizobium meliloti by
                          calorimetric and fluorescence analysis
                          Andujar-Sanchez, Montserrat; Martinez-Rodriguez,
AUTHOR (S):
                          Sergio; Las Heras-Vazguez, Francisco Javier;
                          Clemente-Jimenez, Josefa Maria; Rodriguez-Vico,
                          Felipe; Jara-Perez, Vicente
                          Dpto. Quimica Fisica, Bioquimica y Quimica Inorganica,
CORPORATE SOURCE:
                          Universidad de Almeria, Almeria, 04120, Spain
                          Biochimica et Biophysica Acta, Proteins and Proteomics
SOURCE:
                          (2006), 1764(2), 292-298
                          CODEN: BBAPBW; ISSN: 1570-9639
PUBLISHER:
                          Elsevier B.V.
DOCUMENT TYPE:
                          Journal
                          English
LANGUAGE:
     Hydantoin racemase enzyme together with a
     stereoselective hydantoinase and a stereospecific D-carbamoylase guarantee
     the total conversion from D,L-5-monosubstituted hydantoins with a low
     velocity of racemization, to optically pure D-amino acids.
     Hydantoin racemase from Sinorhizobium meliloti was
     expressed in Escherichia coli. Calorimetric and fluorescence expts. were
     then carried out to obtain the thermodn. binding parameters, \Delta G,
     \Delta H and \Delta S for the inhibitors L- and D-5-methylthioethyl-
     hydantoin. The number of active sites is four per enzyme mol. (one per
     monomer), and the binding of the inhibitor is entropically and
     enthalpically favored under the exptl. conditions studied. In order to
     obtain_information_about_amino_acids_involved_in_the_active_site, four
     different mutants were developed in which cysteines 76 and 181
     were mutated to Alanine and Serine. Their behavior shows that these
     cysteines are essential for enzyme activity, but only cysteine 76 affects
     the binding to these inhibitors.
                                THERE ARE 37 CITED REFERENCES AVAILABLE FOR THIS
REFERENCE COUNT:
                          37
                                RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
     ANSWER 4 OF 9 CAPLUS COPYRIGHT 2007 ACS on STN
                          2004:1127509 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                          142:70788
                          Screening for improved hydantoin
TITLE:
                          racemase mutants, characterization
                          of hydantoin racemase
                          mutants from Arthrobacter crystallopoietes,
                          and their use in the preparation of amino acids
                          May, Oliver; Drauz, Karlheinz; Buchholz, Stefan
INVENTOR(S):
                          Degussa Ag, Germany
PATENT ASSIGNEE(S):
                          PCT Int. Appl., 59 pp.
SOURCE:
                          CODEN: PIXXD2
                                            postdated
DOCUMENT TYPE:
                          Patent
                          English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
                                 DATE
     PATENT NO.
                          KIND
                                              APPLICATION NO.
                                                                      DATE
                          - - - -
                                                                      20040515
     WO 2004111227
                           A2
                                 20041223
                                              WO 2004-EP5239
     WO 2004111227
                                 20050616
                           Α3
         W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH,
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             GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NA, NI, NO,
             NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ,
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TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW
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               AZ, BY, KG, KZ, MD, RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK,
               EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PL, PT, RO, SE,
               SI, SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE,
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     DE 102004022065
                                     20050317
                                                   DE 2004-102004022065
                              A1
                                                   US 2005-559434
     US 2006210989
                              A1
                                     20060921
PRIORITY APPLN. INFO.:
                                                   DE 2003-10326109
                                                                          A 20030606
                                                   DE 2004-102004022065A 20040505
                                                   WO 2004-EP5239
                                                                         W 20040515
      The present invention relates to a screening process for hydantoin
AB
      racemases and to novel hydantoin racemases, to
      the nucleic acid sequences coding therefor and to a process for
     mutagenesis. Hydantoin racemases are of interest in
     connection with the production of enantiomerically enriched amino acids and
     N-carbamoyl amino acids from racemic hydantoins. More specifically,
     preparation of mutant hydantoin racemases from
     Arthrobacter crystallopoietes with improved enzymic properties is
     described. The nucleotide sequences and the encoded amino acid sequences
     of the improved mutant hydantoin racemases
      from A. crystallopoietes are disclosed.
     ANSWER 5 OF 9 CAPLUS COPYRIGHT 2007 ACS on STN DUPLICATE 4
                             2000:707276 CAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                             133:278038
TITLE:
                             Hydantoinase variants with improved
                             properties and their use for the production of amino
                             acids
                             Arnold, Frances H.; May, Oliver; Drauz, Karlheinz;
INVENTOR(S):
                             Bommarius, Andreas
PATENT ASSIGNEE(S):
                             California Institute of Technology, USA; Degussa-Huls
                             A.-G.
SOURCE:
                             PCT Int. Appl., 50 pp.
                             CODEN: PIXXD2
DOCUMENT TYPE:
                             Patent
LANGUAGE:
                             English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
                                    __DATE___
      PATENT NO.
                             KIND
                                                 APPLICATION NO.
                                                                              DATE
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                                     20001005
                                                )WO 2000-US8159
     WO 2000058449
         2000058449

W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, GM, HR, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZW

RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
                             A1
                                                                              20000328
     US 6524837
                              B1
                                     20030225
                                                 US 2000-497585
                                                                              20000203
     EP 1165763
                              A1
                                     20020102
                                                   EP 2000-921477
                                                                              20000328
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                                     20050622
                              B1
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               IE, SI, LT, LV, FI, RO
      JP 2003521235
                              Т
                                     20030715
                                                   JP 2000-608730
                                                                              20000328
     AT 298365
                              Т
                                     20050715
                                                   AT 2000-921477
                                                                              20000328
     EP 1586636
                             A1
                                     20051019
                                                   EP 2004-26624
                                                                              20000328
              AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
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20051031

20051216

PT 2000-921477

ES 2000-921477

US 1999-126923P

20000328

20000328

P 19990329

IE, FI, CY

T3

PT 1165763

ES 2244430

PRIORITY APPLN. INFO.:

US 1999-157427P P 19991004 US 2000-497585 A 20000203 A3 20000328 EP 2000-921477 WO 2000-US8159 W 20000328

Hydantoinase enzymes which are mutants of Arthrobacter DSM-9771 AB hydantoinase are disclosed. The mutants include amino acid substitutions at positions 95, 154, 180, 251 and/or 255 of the wild type hydantoinase. The mutant hydantoinases, like the parent hydantoinase, are used in the production of optically pure amino acids. a mutant gene encoding Arthrobacter hydantoinase with 195L, V180A, and Q251R substitutions was prepared This gene, as well as an Arthrobacter carbamoylase gene, was expressed in Escherichia coli. When used to prepare L-Met from D,L-Met hydantoin, the recombinant bacteria produced about 65 mM L-Met per h while the bacteria expressing the wild-type hydantoinase produced only 8 mM per h. The mutant enzyme was not enantioselective, but was 4-fold more active than the wild-type enzyme.

REFERENCE COUNT:

THERE ARE 9 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 6 OF 9 CAPLUS COPYRIGHT 2007 ACS on STN DUPLICATE 5

ACCESSION NUMBER: 2000:188897 CAPLUS

DOCUMENT NUMBER: 132:278227

TITLE: Inverting enantioselectivity by directed evolution of

hydantoinase for improved production of L-methionine

AUTHOR (S): May, Oliver; Nguyen, Peter T.; Arnold, Frances H.

Division of Chemistry and Chemical Engineering, CORPORATE SOURCE:

California Institute of Technology, Pasadena, CA,

91125, USA

SOURCE: Nature Biotechnology (2000), 18(3), 317-320

CODEN: NABIF9; ISSN: 1087-0156

PUBLISHER: Nature America

DOCUMENT TYPE: Journal LANGUAGE: English

Using directed evolution, the hydantoinase process for production of L-methionine (L-met) in Escherichia coli was improved. This was accomplished by inverting the enantioselectivity and increasing the total activity of a key enzyme in a whole-cell catalyst. The selectivity of all known hydantoinases for D-5-(2-methylthioethyl) hydantoin (D-MTEH) over the L-enantiomer leads to the accumulation of intermediates and reduced productivity for the L-amino acid. Random mutagenesis, saturation mutagenesis, and screening were used to convert the D-selective hydantoinase from Arthrobacter sp. DSM 9771 into an L-selective enzyme and increased its total activity 5-fold. Whole E. coli cells expressing the evolved L-hydantoinase, an L-N-carbamoylase, and a hydantoin racemase produced 91 mM L-met from 100 mM D,L-MTEH in less than 2 The improved hydantoinase increased productivity fivefold for >90% conversion of the substrate. The accumulation of the unwanted intermediate D-carbamoyl-methionine was reduced fourfold compared to cells with the wild-type pathway. Highly D-selective hydantoinase mutants were also discovered. Enantioselective enzymes rapidly optimized by directed evolution and introduced into multienzyme pathways may lead to improved whole-cell catalysts for efficient production of chiral compds.

REFERENCE COUNT:

THERE ARE 22 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 7 OF 9 CAPLUS COPYRIGHT 2007 ACS on STN

22

ACCESSION NUMBER: 2000:327334 CAPLUS

TITLE: Inverting enantioselectivity of a key enzyme creates a

viable process for production of L-methionine.

AUTHOR (S):

Arnold, Frances H.; Nguyen, Peter T.; May, Oliver CORPORATE SOURCE: Division of Chemistry and Chemical Engineering, MC 210-41, California Institute of Technology, Pasadena, CA, 91125, USA

SOURCE:

Book of Abstracts, 219th ACS National Meeting, San Francisco, CA, March 26-30, 2000 (2000), BIOT-112.

American Chemical Society: Washington, D. C.

CODEN: 69CLAC

DOCUMENT TYPE:

Conference; Meeting Abstract

LANGUAGE: English

We have dramatically improved the hydantoinase process for production of L-methionine (L-met) in E. coli by inverting the enantioselectivity of a key enzyme using directed evolution. All known hydantoinases are selective for D-5-(2-methylthioethyl)hydantoin (D-MTEH) over the L-enantiomer, which leads to the accumulation of intermediates and reduced productivity for the L-amino acid. Using random mutagenesis, saturation mutagenesis and screening, we converted the D-selective hydantoinase from Arthrobacter sp. DSM 9771 (eeD=40% at 30% conversion) into an L-selective enzyme (eeL=20%) and increased its catalytic activity 5-fold. coli cells expressing the evolved L-hydantoinase, an L-N-carbamoylase, and a hydantoin racemase produced 91 mM L-met from 100 mM D, L-MTEH in less than 2 h. The improved hydantoinase increased productivity 5-fold for > 90% conversion of the substrate. The accumulation of the unwanted intermediate D-carbamoyl-methionine was reduced 4-fold compared to cells with the wild-type pathway. Highly D-selective (<90%eeD at 30% conversion) hydantoinase mutants were also discovered. Enantioselective enzymes rapidly optimized by directed evolution and introduced into multi-enzyme pathways lead to improved whole-cell catalysts for efficient production of chiral compds.

L4 ANSWER 8 OF 9 BIOTECHDS COPYRIGHT 2007 THE THOMSON CORP. on STN

DUPLICATE 6

ACCESSION NUMBER: 1996-12962 BIOTECHDS

TITLE:

AUTHOR:

New microorganism DSM 9771 and use for recovery of gene coded

with carbamoylase, hydantoinase or hydantoin-

racemase;

L-alpha-amino acid e.g. methionine production Wagner F; Hantke B; Wagner T; Drauz K; Bommarius A

PATENT ASSIGNEE: Degussa

LOCATION: Frankfurt, Germany.

PATENT INFO: DE 19519717 22 Aug 1996

APPLICATION INFO: DE 1995-119717 30 May 1995

PRIORITY INFO: DE 1995-119717 30 May 1995

DOCUMENT TYPE: Patent LANGUAGE: German

OTHER SOURCE: WPI: 1996-372118 [38]

A microorganism, DSM 9771, (or a mutant) harboring genes encoding carbamoylase (I), and/or dihydropyrimidinase (EC-3.5.2.2), or hydantoin-racemase, is claimed. DSM 9771 (or enzymes from the microorganism) can be used to produce L-alpha-amino acids, by enzymatic conversion of a D-, L-, and/or D,L,N-carbamoyl-alpha-amino acid. The new microorganism has high activity and produces a series of enzymes in large amounts. In an example, a wild strain of DSM 7330 was cultured 10 times on 5 ml of a culture medium containing 10.0 g/lglucose, 0.95 g/l KH2PO4, 2.0 g/l K2HPO4.3H2O, 0.2 g/l MgSO4.7H2O, 0.02 g/l CaCl2.2H2O, 2.0 g/l L-carbamoylmethionine, and 10 ml/l of a solution of trace elements. Single colonies were isolated and incubated on agar plates for 4 days at 30 deg, and then stored in a culture medium containing 10.0 g/l yeast extract, 10.0 g/l bactopeptone, 10.0 g/l glucose, 3.0 g/l NaCl, 0.1 g/l MgCl2.4H2O, 10 ml/l trace solution, and 15.0 g/l agar. Under N2, a suspension of 15 g of DSM 9771 and 2.25 g of D,L-methylthioethylhydantoin in 150 ml of saline was incubated at 37 deg and pH 8.5 for 30 hr. Separation yielded 1.6 g (82.6%) of L-methionine. (10pp)

ACCESSION NUMBER: 1995-01724 BIOTECHDS

TITLE: New microorganism for L-alpha-amino acid production;

dihydropyrimidinase and hydantion-racemase gene for use in

strain improvement for monosubstituted hydantoin or

carbamoyl-alpha-amino acid conversion

AUTHOR: Wagner F; Voelkel D; Bommarius A; Drauz K

PATENT ASSIGNEE: Degussa

PATENT INFO: EP 625571 23 Nov 1994
APPLICATION INFO: EP 1994-107323-11 May 1994
PRIORITY INFO: DE 1993-4316928 19 May 1993

DOCUMENT TYPE: Patent LANGUAGE: German

OTHER SOURCE: WPI: 1995-359746 [01]

AB New microorganisms DSM 7329 and DSM 7330 are claimed for use in L-alpha-amino acid production from D-, L- and/or D,L-5-monosubstituted hydantoins and/or D,L-N-carbamoyl-alpha-amino acids. Also claimed are mutants and variants of the new microorganisms; use of

the mutants, variants and isolated enzymes

(dihydropyrimidinase (EC-3.5.2.2) and/or hydantion-racemase) from the microorganisms for L-amino acid production; and the enzyme encoding genes from the new microorganisms for use in cell transformation to facilitate L-amino acid production. (18pp)

# **Refine Search**

### Search Results -

Term	Documents
HYDANTOIN	9967
HYDANTOINS	3953
RACEMASE	3575
RACEMASES	372
(HYDANTOIN ADJ RACEMASE).CLMPGPB,USPT,USOC,EPAB,JPAB,DWPI.	19
((HYDANTOIN RACEMASE).CLM. ).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	19

US Pre-Grant Publication Full-Text Database US Patents Full-Text Database

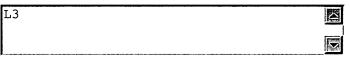
US OCR Full-Text Database

Database:

EPO Abstracts Database
JPO Abstracts Database
Derwent World Patents Index

**IBM Technical Disclosure Bulletins** 

Search:











### **Search History**

DATE: Tuesday, February 27, 2007 Purge Queries Printable Copy Create Case

Set Name side by side	Query	Hit Count	Set Name result set
•	LICAT LICAC ED AD IDAD DIVIDI. DI LID—	VEC. OD- 4D I	
DB=PGPB,	USPT, USOC, EPAB, JPAB, DWPI; PLUR =	IES; OP=ADJ	
<u>L3</u>	(hydantoin racemase).clm.	19	<u>L3</u>
<u>L2</u>	L1 and (position 79)	2	<u>L2</u>
<u>L1</u>	hydantoin racemase	. 70	<u>L1</u>

**END OF SEARCH HISTORY** 

## **Hit List**

First Hit Clear Generate Collection Print Fwd Refs Bkwd Refs

Generate OACS

**Search Results -** Record(s) 1 through 2 of 2 returned.

☐ 1. Document ID: US 20060210989 A1

L2: Entry 1 of 2

File: PGPB

Sep 21, 2006

PGPUB-DOCUMENT-NUMBER: 20060210989

PGPUB-FILING-TYPE:

DOCUMENT-IDENTIFIER: US 20060210989 A1

TITLE: Screening process for hydantoin racemases

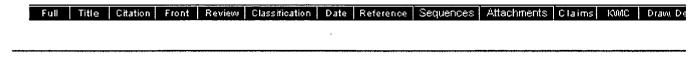
PUBLICATION-DATE: September 21, 2006

INVENTOR-INFORMATION:

NAME CITY STATE COUNTRY

May; Oliver Frankfurt DE
Drauz; Karlheinz Freigericht DE
Buchholz; Stefan Hanau DE

US-CL-CURRENT: <u>435/6</u>; <u>435/106</u>, <u>435/15</u>, <u>435/233</u>, <u>435/325</u>, <u>536/23.2</u>



2. Document ID: US 20060210989 A1, WO 2004111227 A2, DE 1004022065 A1

L2: Entry 2 of 2

File: DWPI

Sep 21, 2006

DERWENT-ACC-NO: 2005-066238

DERWENT-WEEK: 200663

COPYRIGHT 2007 DERWENT INFORMATION LTD

TITLE: Screening for  $\underline{\text{hydantoin racemases}}$  by allowing enantioselective hydantoinase and  $\underline{\text{hydantoin racemase}}$  to act on chiral hydantoin and detecting obtained N-carbamoylamino acid or freed protons

Full Title Citation Front Review Classification Date Reference Table 100 Claims KWC Draw De Clear Generate Collection Print Fwd Refs Bkwd Refs Generate OACS

Term Documents POSITION 6741003

POSITIONS	2338275
"79"	900142
79S	213
(1 AND (POSITION ADJ "79")).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	2
(L1 AND (POSITION 79) ).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	2

Display Format:	_	Change Format

Previous Page Next Page Go to Doc#

# **Hit List**

First Hit Clear Generate Collection Print Fwd Refs Bkwd Refs
Generate OACS

Search Results - Record(s) 1 through 19 of 19 returned.

☐ 1. Document ID: US 20060246553 A1

L3: Entry 1 of 19

File: PGPB

Nov 2, 2006

PGPUB-DOCUMENT-NUMBER: 20060246553

PGPUB-FILING-TYPE:

DOCUMENT-IDENTIFIER: US 20060246553 A1

TITLE: 5-substituted hydantoin racemase, DNA coding for the racemase, and processes

for producing optically active amino acids

PUBLICATION-DATE: November 2, 2006

INVENTOR-INFORMATION: .

NAME CITY STATE COUNTRY

Suzuki; Shunichi Kanagawa JP
Onishi; Norimasa Kanagawa JP
Yokozeki; Kenzo Kanagawa JP

US-CL-CURRENT: 435/108

Full Ti	itle   Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWC	Draw, De
				***							

☐ 2. Document ID: US 20060210989 A1

L3: Entry 2 of 19

File: PGPB

Sep 21, 2006

PGPUB-DOCUMENT-NUMBER: 20060210989

PGPUB-FILING-TYPE:

DOCUMENT-IDENTIFIER: US 20060210989 A1

TITLE: Screening process for hydantoin racemases

PUBLICATION-DATE: September 21, 2006

INVENTOR-INFORMATION:

NAME CITY STATE COUNTRY

May; Oliver Frankfurt DE
Drauz; Karlheinz Freigericht DE
Buchholz; Stefan Hanau DE

US-CL-CURRENT: 435/6; 435/106, 435/15, 435/233, 435/325, 536/23.2

Page 2 of 9

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWIC	Draw. De
												<del></del>

### ☐ 3. Document ID: US 20060035321 A1

L3: Entry 3 of 19

File: PGPB

Feb 16, 2006

Nov 3, 2005

PGPUB-DOCUMENT-NUMBER: 20060035321

PGPUB-FILING-TYPE:

DOCUMENT-IDENTIFIER: US 20060035321 A1

TITLE: Hydantoin racemase

PUBLICATION-DATE: February 16, 2006

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Boesten; Wilhelmus Hubertus Joseph	Sittard		NL
Kierkels; Joannes Gerardus Theodorus	Sittard		NL
Assema; Bernard Jan	Geleen		NL
Fuiz Perez; Luis Miguel	Granada		ES
Gonzalez Pacanowska; Dolores	Granada		ES
Gonzalez Lopez; Jesus	Granada		ES
De La Escal Era Hueso; Santiago	Mojaca		ES

US-CL-CURRENT: 435/69.1; 435/106, 435/233, 435/320.1, 435/325, 536/23.2

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWC	Draw, De
											***************************************	
	4.	Documen	nt ID:	US 20	050244936	<b>A</b> 1						

File: PGPB

PGPUB-DOCUMENT-NUMBER: 20050244936

PGPUB-FILING-TYPE: new

L3: Entry 4 of 19

DOCUMENT-IDENTIFIER: US 20050244936 A1

TITLE: Hydantoin-racemase

PUBLICATION-DATE: November 3, 2005

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Altenbuchner, Josef	Nufringen	DE	DE
Mattes, Ralf	Stuttgart	GA	DE
Pietzsch, Markus	Halle		DE
Syldatk, Christoph	Stuttgart		US
Wiese, Anja	Eching		DE
Bommarius, Andreas	Atlanta		US

Record List Display Page 3 of 9

Tischer, Wilhelm

Peissenberg

DE

US-CL-CURRENT: 435/106; 435/233, 435/252.3, 435/471, 536/23.2, 536/24.3

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw. De

☐ 5. Document ID: US 20050214912 A1

L3: Entry 5 of 19

File: PGPB

Sep 29, 2005

PGPUB-DOCUMENT-NUMBER: 20050214912

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20050214912 A1

TITLE: Method for producing an optically active amino acid

PUBLICATION-DATE: September 29, 2005

INVENTOR-INFORMATION:

NAME CITY STATE COUNTRY

Nozaki, Hiroyuki Kawasaki-shi JP Watanabe, Kunihiko Kawasaki-shi JP

US-CL-CURRENT: 435/106; 435/233, 435/252.33

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw De

☐ 6. Document ID: US 20050202542 A1

L3: Entry 6 of 19

File: PGPB

Sep 15, 2005

PGPUB-DOCUMENT-NUMBER: 20050202542

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20050202542 A1

TITLE: Production method of D-alloisoleucine

PUBLICATION-DATE: September 15, 2005

INVENTOR-INFORMATION:

NAME CITY STATE COUNTRY

Maruyama, Shogo Kawasaki-shi JP Kira, Ikuo Kawasaki-shi JP Takemoto, Tadashi Kawasaki-shi JP

US-CL-CURRENT: 435/106

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw. De

7. Document ID: US 20050112729 A1

L3: Entry 7 of 19

File: PGPB

May 26, 2005

Apr 21, 2005

Sep 18, 2003

PGPUB-DOCUMENT-NUMBER: 20050112729

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20050112729 A1

TITLE: Recombinant DNA having hydantoinase gene and carbamylase gene and process

for producing amino acid

PUBLICATION-DATE: May 26, 2005

INVENTOR-INFORMATION:

NAME CITY STATE COUNTRY JP Kira, Ikuo Kanagawa JP Takenaka, Yasuhiro Kanagawa JΡ Nozaki, Hiroyuki Kanagawa JΡ Watanabe, Kunihiko Kanagawa

US-CL-CURRENT: 435/106; 435/228, 435/252.33

	***********
☐ 8. Document ID: US 20050084946 A1	

File: PGPB

PGPUB-DOCUMENT-NUMBER: 20050084946

PGPUB-FILING-TYPE: new

L3: Entry 8 of 19

DOCUMENT-IDENTIFIER: US 20050084946 A1

TITLE: 5-substituted hydantoin racemase, DNA coding for the racemase, and processes

for producing optically active amino acids

PUBLICATION-DATE: April 21, 2005

INVENTOR-INFORMATION:

L3: Entry 9 of 19

NAME CITY STATE COUNTRY
Suzuki, Shunichi Kanagawa JP
Onishi, Norimasa Kanagawa JP
Yokozeki, Kenzo Kanagawa JP

US-CL-CURRENT: 435/233; 435/252.3, 435/320.1, 435/6, 435/69.1, 536/23.2

Full Title	e Citation Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWC	Draw De
								•		
							·		4-3-4	
□ 9.	Document ID:	US 20	030175910	A1						

File: PGPB

Record List Display Page 5 of 9

PGPUB-DOCUMENT-NUMBER: 20030175910

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20030175910 A1

TITLE: Whole cell catalyst

PUBLICATION-DATE: September 18, 2003

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Altenbuchner, Josef	Nufringen	GA	DE
Bommarius, Andreas	Atlanta		US
Mattes, Ralf	Stuttgart		DE
Syldatk, Christoph	Stuttgart		DE
Tischer, Wilhelm	Peissenberg		DE
Wiese, Anja	Eching		DE
Wilms, Burkard	Stuttgart		DE

US-CL-CURRENT: 435/106; 435/228, 435/252.33

Full Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWWC	Draw, De
□ 10	Dogume	ant ID	. 110 2	003016617	Q Λ 1				.,		
I 10.	Docume	בווו זוט	. 032	003010017	o Ai			•			
L3: Entry	10 of	19				File:	PGPB		Sep	4,	2003

PGPUB-DOCUMENT-NUMBER: 20030166178

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20030166178 A1

TITLE: Process for the production of enantiomer-enriched alpha-substituted

carboxylic acids

PUBLICATION-DATE: September 4, 2003

INVENTOR-INFORMATION:

NAME .	CITY	STATE	COUNTRY
May, Oliver	Frankfurt		DE
Syldatk, Christoph	Stuttgart		DE
Vielhauer, Oliver	Stuttgart		DE
Werner, Markus	Weinsberg		DE

US-CL-CURRENT: <u>435/136</u>

Full Title (	Citation Front	Review	Classification	Date	Referenc	se Sequences	Attachments	Claims	KMC	Draw De
□ 11. I	Document ID	: US 2	003014324	4 A1						
L3: Entry 1	.1 of 19				File:	PGPB		Jul	31,	2003

Record List Display Page 6 of 9

PGPUB-DOCUMENT-NUMBER: 20030143244

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20030143244 A1

TITLE: Arthrobacter D-carbamoylase and methods of preparing enantiomerically

enriched D-amino acids

PUBLICATION-DATE: July 31, 2003

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Drauz, Karlheinz	Freigericht	GA	DE
May, Oliver	Frankfurt		DE
Bommarius, Andreas	Atlanta		US
Syldatk, Christoph	Stuttgart		DE .
Altenbuchner, Josef	Nufringen		DE
Werner, Markus	Weinsberg		DE
Siemann-Herzberg, Martin	Wildberg		DE

US-CL-CURRENT: 424/190.1; 435/196, 435/252.3, 435/320.1, 435/69.3, 536/23.2

Full	Title	Citation Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWIC	Draw, De
 				·						<del></del>	
	12.	Document ID	: US 2	002013284	8 A 1						
*		200000000000000000000000000000000000000									

File: PGPB

PGPUB-DOCUMENT-NUMBER: 20020132848

PGPUB-FILING-TYPE: new

L3: Entry 12 of 19

DOCUMENT-IDENTIFIER: US 20020132848 A1

TITLE: Process for the preparation of allysine acetal

PUBLICATION-DATE: September 19, 2002

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY
Krimmer, Hans-Peter	Dietzenbach		DE -
May, Oliver	Frankfurt		DE
Klement, Ingo	Pohlheim-Garbenteich		DE
Drauz, Karlheinz	Freigericht		DE
Reichert, Dietmar	Eschau		DE

US-CL-CURRENT: <u>514/561</u>; <u>435/106</u>

_ Full Title_	Citation From	nt Review	Classification	Date	Reference	Sequences	Attachments	Claims	KMC	Draw, De
					<u> </u>					

### ☐ 13. Document ID: US 20020102713 A1

Sep 19, 2002

Page 7 of 9 Record List Display

L3: Entry 13 of 19

File: PGPB

Aug 1, 2002

PGPUB-DOCUMENT-NUMBER: 20020102713

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020102713 A1

TITLE: 5-substituted hydantoin racemase, DNA coding for the racemase, and processes

for producing optically active amino acids

PUBLICATION-DATE: August 1, 2002

INVENTOR-INFORMATION:

CITY STATE COUNTRY NAME

Suzuki, Shunichi Kanagawa JΡ Onishi, Norimasa JP Kanagawa JP Yokozeki, Kenzo Kanagawa

US-CL-CURRENT: 435/233; 435/106, 435/320.1, 435/325, 435/69.1, 536/23.2

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw. De ☐ 14. Document ID: US 6825014 B2

L3: Entry 14 of 19

File: USPT

Nov 30, 2004

US-PAT-NO: 6825014

DOCUMENT-IDENTIFIER: US 6825014 B2

TITLE: Process for the preparation of allysine acetal

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw De ☐ 15. Document ID: US 6815195 B2 L3: Entry 15 of 19 Nov 9, 2004 File: USPT

US-PAT-NO: 6815195

DOCUMENT-IDENTIFIER: US 6815195 B2

TITLE: 5-substituted hydantoin racemase, DNA coding for the racemase, and processes

for producing optically active amino acids

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw De

☐ 16. Document ID: US 6800464 B2

L3: Entry 16 of 19

File: USPT

Oct 5, 2004

US-PAT-NO: 6800464

DOCUMENT-IDENTIFIER: US 6800464 B2

Record List Display Page 8 of 9

TITLE: Arthrobacter D-carbamoylase and methods of preparing enantiomerically enriched D-amino acids

Full Title Citation Front Review Classification Date Reference Sequences Attachinade Claims KMC Draw De 17. Document ID: US 6713288 B1
L3: Entry 17 of 19 File: USPT Mar 30, 2004

US-PAT-NO: 6713288

DOCUMENT-IDENTIFIER: US 6713288 B1

\*\* See image for Certificate of Correction \*\*

TITLE: Whole cell catalysts

Full Title Citation Front Review Classification Date Reference Sequences Stackments Claims KWC Draw De

18. Document ID: US 5827717 A

L3: Entry 18 of 19 File: USPT Oct 27, 1998

US-PAT-NO: 5827717

DOCUMENT-IDENTIFIER: US 5827717 A

TITLE: Microorganisms their use and method of producing L-.alpha.-amino acids

Full Title Citation Front Review Classification Date Reference Sequence Statements Claims KWC Draw De ...

19. Document ID: US 5516660 A

L3: Entry 19 of 19 File: USPT May 14, 1996

US-PAT-NO: 5516660

DOCUMENT-IDENTIFIER: US 5516660 A

TITLE: Microorganisms, their use and method of producing L-.alpha.-amino acids

Generate Collection Print Fwd F	Refs Bkwd Refs Generate
Term	Documents
HYDANTOIN	9967
HYDANTOINS	. 3953
RACEMASE	3575
RACEMASES	372

Full Title Citation Front Review Classification Date Reference Settlements Claims Killic Draw De

(HYDANTOIN ADJ RACEMASE).CLMPGPB,USPT,USOC,EPAB,JPAB,DWPI.	19
((HYDANTOIN RACEMASE).CLM. ).PGPB,USPT,USOC,EPAB,JPAB,DWPI.	19

Display Format:  -	Change Format
--------------------	---------------

Previous Page Next Page Go to Doc#

```
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Q982Y4_RHILO-
                                                          All Wele are
G797 D
Risoltifution
ID
    Q982Y4 RHILO
                   PRELIMINARY;
                                 PRT;
                                        250 AA.
AC
    Q982Y4;
DT
     01-OCT-2001, integrated into UniProtKB/TrEMBL.
DT
     01-OCT-2001, sequence version 1.
DT
     07-FEB-2006, entry version 16.
DΕ
    Hydantoin racemase.
GN
    OrderedLocusNames=mll8441;
os
    Rhizobium loti (Mesorhizobium loti).
OC
    Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
OC
    Phyllobacteriaceae; Mesorhizobium.
OX
    NCBI TaxID=381;
RN
RP
    NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RC
    STRAIN=MAFF303099;
    MEDLINE=21082930; PubMed=11214968; DOI=10.1093/dnares/7.6.331;
RX
    Kaneko T., Nakamura Y., Sato S., Asamizu E., Kato T., Sasamoto S.,
RA
    Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T.,
RA
    Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno A.,
RA
    Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto M.,
RA
RA
    Takeuchi C., Yamada M., Tabata S.;
RT
    "Complete genome structure of the nitrogen-fixing symbiotic bacterium
RT
    Mesorhizobium loti.";
RL
    DNA Res. 7:331-338(2000).
CC
CC
    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
CC
    Distributed under the Creative Commons Attribution-NoDerivs License
CC
    EMBL; BA000012; BAB54322.1; -; Genomic DNA.
DR
DR
    BioCyc; MLOT381:MLL8441-MONOMER; -.
DR
    GO; GO:0016855; F:racemase and epimerase activity, acting on . . .; IEA.
DR
    GO; GO:0008152; P:metabolism; IEA.
DR
    InterPro; IPR001920; Asp/Glu race.
DR
    Pfam; PF01177; Asp_Glu_race; 1.
KW
    Complete proteome.
SQ
    SEQUENCE
               250 AA; 25707 MW; CFBE8893D6A762F8 CRC64;
 Query Match
                        39.8%;
                                Score 466; DB 2;
                                                  Length 250;
 Best Local Similarity
                        43.0%;
                                Pred. No. 1.1e-26;
 Matches 101; Conservative
                              44; Mismatches
                                                                         2;
                                              88;
                                                   Indels
                                                             2;
                                                                 Gaps
Qу
           1 MRILVINPNSSSALTESVADAAQQVVATGTIISAINPSRGPAVIEGSFDEALATFHLIEE 60
                                       ::|||:|||:||
Db
           3 VQILVVNPNTTASMTETIGAAARAVAGAWTEIIAVTSSTGPDSIEGYYDEALAVPGLLME 62
          61 VERAERENPPDAYVIACFGDPGLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPR 120
Qy
                        Db
          63 IAAGERRG-AQAAVIACFDDTGLDAARAMANIPVIGICEAALSMASFIAQRFTVVTTTER 121
         121 VRKHLHELVRQAGATNRLASIKLPNLGVMAFHEDEHAALETLKQAAKEAVQEDGAESIVL 180
Qу
              |:
Db
         122 SRVPVEGLVQRYGMAGR-ARVRAADIPVLALEDPASGATGKLRDEIARAVEEDRAEAIVL 180
Qy
         181 GCAGMVGFARQLSDELGVPVIDPVEAACRVAESLVALGYQTSKANSYQKPTEKQY 235
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ΙD
XX
AC
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XX
DT
     11-MAR-2004 (first entry)
XX
DE
     Agrobacterium radiobacter hydantoin racemase.
XX
KW
     Hydantoin racemase; enzyme.
XX
os
     Agrobacterium tumefaciens.
XX
PN
     WO2003100050-A1.
XX
PD
     04-DEC-2003.
XX
     23-MAY-2003; 2003WO-NL000386.
PF
XX
PR
     23-MAY-2002; 2002NL-01020663.
XX
PA
     (STAM ) DSM NV.
XX
     Boesten WHJ, Kierkels JGT, Assema FBJ, Ruiz Perez LM;
PΙ
PΙ
     Gonzalez Pacanowska D, Gonzalez Lopez J, De La Escalera Hueso S;
XΧ
DR
     WPI; 2004-081908/08.
DR
     N-PSDB; ADG73872.
XX
PT
     New polypeptide with hydantoin racemase activity, that does not suffer
PT
     from substrate inhibition, useful for catalyzing the racemization of
PT
     substituted D or L hydantoins.
XX
PS
     Claim 2; SEQ ID NO 2; 27pp; English.
XX
CC
     The present sequence is the protein sequence of Agrobacterium radiobacter
CC
     hydantoin racemase, an enzyme that catalyses the racemisation of
CC
     substituted D or L hydantoins. The A. radiobacter enzyme differs from
CC
     other hydantoin racemases as it does not exhibit substrate inhibition.
CC
     The invention provides processes for the racemisation of enantiomerically
CC
     enriched hydantoin compounds and to processes for the preparation of
     enantiomerically enriched D- or L-alpha amino acids using the hydantoin
CC
CC
     racemase. Examples from the invention describe the expression of A.
CC
     radiobacter hydantoin racemase in Escherichia coli, and use of the over-
CC
     expressed enzyme in racemisation experiments using D-5-benzylhydantoin, L
CC
     -5-isopropylhydantoin and L-5-methylmercaptoethylhydantoin substrates.
XX
SQ
     Sequence 238 AA;
 Query Match
                          40.7%;
                                  Score 476.5;
                                                 DB 8;
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 Best Local Similarity
                          44.0%; Pred. No. 4.6e-42;
 Matches 103; Conservative
                                39; Mismatches
                                                   91;
                                                        Indels
                                                                      Gaps
                                                                              1;
                                                                  1:
Qу
            1 MRILVINPNSSSALTESVADAAQQVVATGTIISAINPSRGPAVIEGSFDEALATFHLIEE 60
              1 1 :11111::::1
                                1:1:1
                                          1 111 11
                                                       1 111 11111
Db
            1 MHIRLINPNSTASMTAQALDSALRVKQAHTTISATNPLDTPVSIEGGADEALAVPGMLEE 60
Qy
           61 VERAERENPPDAYVIACĖGDPGLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPR 120
              : : 11
                        11111111 1111 1 :1:
                                               1|:|: :| | :: :: |||:: |||
Db
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Qy	181	GCAGMVGFARQLSDELGVPVIDPVEAACRVAESLVALGYQTSKANSYQKPTEKQ 234
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EndFr</td <td>agme</td> <td>nt&gt;</td>	agme	nt>

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<!--StartFragment-->RESULT 18
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ID
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AC
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XX
DT
    26-FEB-2004 (first entry)
XX
DE
    Flavobacterium hydantoin racemase amino acid sequence.
XX
KW
    hydantoin racemase; Flavobacterium;
KW
    optically active amino acid production; enzyme.
XX
OS
    Flavobacterium sp.
XX
PN
    JP2003210176-A.
XX
PD
    29-JUL-2003.
XX
    22-JAN-2002; 2002JP-00013552.
PF
XX
PR
    .22-JAN-2002; 2002JP-00013552.
XX
PA
    (AJIN ) AJINOMOTO KK.
XX
DR
    WPI; 2003-819831/77.
DR
    N-PSDB; ADF86382, ADF86392.
XX
PT
    Novel protein having penta substituted hydantoin racemase, activity,
PT
    useful for manufacturing optically active amino acid.
XX
    Claim 1; SEQ ID NO 2; 30pp; Japanese.
PS
XX
    This invention relates to a novel protein having penta substituted
CC
    hydantoin racemase activity (derived from Flavobacterium) which comprises
CC
    a fully defined sequence of 243 amino acids as given in the
CC
CC
    specification, optionally containing deletion, insertion, addition or
    inversion of one or more amino acid residues. The invention is useful in
CC
CC
    the production of optically active amino acids.
XX
SO
    Sequence 243 AA;
 Query Match
                        40.6%; Score 474.5; DB 7;
                                                  Length 243;
 Best Local Similarity
                        41.7%; Pred. No. 7.8e-42;
           98; Conservative
                             45; Mismatches
                                              91;
                                                  Indels
                                                              Gaps
                                                                       1;
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Qу
             Db
           1 MKIKVINPNTTLTMTAKIGEAAAAVASAGTEVVAVSPAMGPASIEGHYDEAVSALGVLDE 60
          61 VERAERENPPDAYVIACFGDPGLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPR 120
Qу
                      Db
          61 VRKGKAEG-CDGYLIACFDDPGLQAAREIADGPVVGIAEAAMHMASFVSEGFSVVATGHR 119
         121 VRKHLHELVRQAGATNRLASIKLPNLGVMAFHEDEHAALETLKQAAKEAVQEDGAESIVL 180
Qу
                Db
         120 SRIILEHLARSYGMEHKCRKVRTTELAVLDLEVEGSDARGIILEECRRAIVEDHSDCIVL 179
Qy
         181 GCAGMVGFARQLSDELGVPVIDPVEAACRVAESLVALGYQTSKANSYQKPTEKQY 235
                   | | | | | | | | |
Dh
         180 GCAGMADLADYISKELGVPVVDGVAAGVKVLEGLIGLRLSTSRACGYAYPNPKTY 234
```

<!--EndFragment-->

```
<!--StartFragment-->RESULT 16
ADG73873-
ID
     ADG73873 standard; protein; 238 AA.
XX
AC
     ADG73873;
XX
DT
     11-MAR-2004 (first entry)
XX
     Agrobacterium radiobacter hydantoin racemase.
DE
XX
     Hydantoin racemase; enzyme.
KW
XX
os
     Agrobacterium tumefaciens.
XX
PN
     WO2003100050-A1.
XX
PD
     04-DEC-2003.
XX
     23-MAY-2003; 2003WO-NL000386.
PF
XX
     23-MAY-2002; 2002NL-01020663.
PR
XX
PA
     (STAM ) DSM NV.
XX
PΙ
     Boesten WHJ, Kierkels JGT, Assema FBJ, Ruiz Perez LM;
     Gonzalez Pacanowska D, Gonzalez Lopez J, De La Escalera Hueso S;
PΙ
XX
     WPI; 2004-081908/08.
DR
     N-PSDB; ADG73872.
DR
XX
PT
     New polypeptide with hydantoin racemase activity, that does not suffer
PT
     from substrate inhibition, useful for catalyzing the racemization of
PT
     substituted D or L hydantoins.
XX
PS
     Claim 2; SEQ ID NO 2; 27pp; English.
XX
CC
     The present sequence is the protein sequence of Agrobacterium radiobacter
CC
     hydantoin racemase, an enzyme that catalyses the racemisation of
CC
     substituted D or L hydantoins. The A. radiobacter enzyme differs from
CC
     other hydantoin racemases as it does not exhibit substrate inhibition.
CC
     The invention provides processes for the racemisation of enantiomerically
CC
     enriched hydantoin compounds and to processes for the preparation of
CC
     enantiomerically enriched D- or L-alpha amino acids using the hydantoin
CC
     racemase. Examples from the invention describe the expression of A.
CC
     radiobacter hydantoin racemase in Escherichia coli, and use of the over-
CC
     expressed enzyme in racemisation experiments using D-5-benzylhydantoin, L
CC
     -5-isopropylhydantoin and L-5-methylmercaptoethylhydantoin substrates.
XX
SQ
     Sequence 238 AA;
  Query Match
                          40.7%;
                                  Score 476.5; DB 8;
                                                        Length 238;
 Best Local Similarity
                          44.0%;
                                  Pred. No. 4.6e-42;
 Matches 103; Conservative
                                39; Mismatches
                                                   91;
                                                        Indels
Qy
            1 MRILVINPNSSSALTESVADAAQQVVATGTIISAINPSRGPAVIEGSFDEALATFHLIEE 60
                                 1:1:1
              1 | :||||::::|
                                           | | | | | |
                                                       1 111 1111
Db
            1 MHIRLINPNSTASMTAQALDSALRVKQAHTTISATNPLDTPVSIEGGADEALAVPGMLEE 60
           61 VERAERENPPDAYVIACFGDPGLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPR 120
Qу
                        11111111 1111 1 :1:
                                               11:1: :1 1 :: :: 111:: 111
Db
           61 IRKGERLG-VDAYVIACFDDPGLHAAREVARGPVIGICQAGIQVAMTISRRFSIITTLPR 119
```

Qy	121	VRKHLHELVRQAGATNRLASIKLPNLGVMAFHEDEHAALETLKQAAKEAVQEDGAESIVL 180
Db	120	::      ::       ::             ::
•		
Qy		GCAGMVGFARQLSDELGVPVIDPVEAACRVAESLVALGYQTSKANSYQKPTEKQ 234
Db	180	GCAGMSALCDRLREATGVPVIDGVTAAVKLAEALVGAGYSTSKVNAYDYPRIKE 233

<!--EndFragment-->

```
<!--StartFragment-->RESULT 9
probable hydantoin racemase - Pyrococcus horikoshii
C; Species: Pyrococcus horikoshii
C; Date: 14-Aug-1998 #sequence_revision 14-Aug-1998 #text change 09-Jul-2004
C; Accession: B71099
R; Kawarabayasi, Y.; Sawada, M.; Horikawa, H.; Haikawa, Y.; Hino, Y.; Yamamoto, S.; Sek
DNA Res. 5, 55-76, 1998
A; Title: Complete sequence and gene organization of the genome of a hyper-thermophilic
A; Reference number: A71000; MUID: 98344137; PMID: 9679194
A; Accession: B71099
A; Status: preliminary; nucleic acid sequence not shown; translation not shown
A; Molecule type: DNA
A; Residues: 1-228 <KAW>
A;Cross-references: UNIPROT:058781; UNIPARC:UPI0000066833; GB:AP000004; NID:g3236131;
A; Experimental source: strain OT3
A; Note: this accession replaces an interim accession for a sequence replaced by GenBan
C; Genetics:
A;Gene: PH1054
                       14.9%; Score 174; DB 2; Length 228;
 Query Match
 Best Local Similarity 28.6%; Pred. No. 1.7e-06;
 Matches 58; Conservative 29; Mismatches 64; Indels 52; Gaps
                                                                      8;
          43 VIEGSFDEALATFHLIEEV-----ERAERENPP------DAYVIACFGDPG 82
Qу
            Db
          29 IIESAFPELKVVSRCIEDQPKGIYNEETEREAEPKIIRLAKEFEREGVDAIIISCAADPA 88
         83 LDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPRVRKHLHELVRQAGATNRLASIK 142
Qу
            Db
          89 VEKVRKLLSIPVIG---AGSSVSALALAYGRRVGVL-NLTEETPKVIRSILGNNLIA--- 141
         143 LPNLGVMAFHEDEHAALET------LKQAAKEAVQEDGAESIVLGCAGM--VGF 188
Qу
                 11 ::
                                        : ||| ::| | | | || :|
         142 -----EDHPSGVSNTLDLLTDWGRREVINAAKR-LKEKGVEVIALGCTGMSTIGI 190
Db
         189 ARQLSDELGVPVIDPVEAACRVA 211
Qу
            1 1:1:1:11111 1: 11
         191 APVLEEEVGIPVIDPVIASGAVA 213
Db
<!--EndFragment-->
```

```
<!--StartFragment-->RESULT 8
T41412
probable hydantoin racemase - fission yeast (Schizosaccharomyces pombe)
C; Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence revision 03-Dec-1999 #text change 09-Jul-2004
C; Accession: T41412
R; Wood, V.; Rajandream, M.A.; Barrell, B.G.; Murphy, L.; Harris, D.
submitted to the EMBL Data Library, September 1998
A; Reference number: Z21954
A; Accession: T41412
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: DNA
A; Residues: 1-236 < WOO>
A; Cross-references: UNIPROT: O74886; UNIPARC: UPI000006AF98; EMBL: AL031798; PIDN: CAA2118
A; Experimental source: strain 972h-; cosmid c576
A; Gene: SPDB: SPCC576.02
A; Map position: 3
 Query Match
                         17.2%; Score 201.5; DB 2;
                                                    Length 236;
 Best Local Similarity
                         29.5%; Pred. No. 1.2e-08;
          69; Conservative 40; Mismatches 118;
                                                     Indels
                                                                 Gaps
           3 ILVINPNSSSALTESVADAAQQVVATGTIISAIN-PSRGPAVIEGSFDEALATFHLIEEV 61
Qу
             4 ILVINPNSSTFITTSMEEKLVPLVPSDVKLRFLTCPQPGAAVID-SITEATLTAALVFQA 62
Db
          62 ERAERENPPDAYVIACFG-DPĞLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPR 120
Qу
                                | :| ::|
                                            | :|: :|:: : |
          63 LTPSVLDGVDAIAVACYSPTPLVDMIRESFALPCMGIVQASVLSALSVGQRIGILTSTYR 122
Db
         121 VRKHLHELVRQAGAT-NRLASIKLPNLGVMAFHEDEHAALET-LKQAAKEAVQEDGAESI 178
Qу
                 1:11: 1:1:1:1
                                        1: :
                                                   | | | | | | | | |
Db
         123 SECLLYELLDSFGVSRTRVAAIASTGRTVLQLSQMPSQERETLLVQKAQELANTKGADVI 182
Qу
         179 VLGCAGMVGFARQLSDELG--VPVIDPVEAACRVAESLVALGYQTSKANSYQKP 230
                         1:
                            111
                                                             1
         183 CLGGAALAAIRDQIQVAVGPNIPIIDGVHAAVELLAGLARQNLHTSKFGIYTYP 236
Db
```

```
<!--StartFragment-->RESULT 5
AD3135
hydantoin racemase [imported] - Agrobacterium tumefaciens (strain C58, Dupont)
C; Species: Agrobacterium tumefaciens
C; Date: 11-Jan-2002 #sequence revision 11-Jan-2002 #text change 09-Jul-2004
C; Accession: AD3135
R; Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo,
Science 294, 2317-2323, 2001
A; Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm
A; Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58.
A; Reference number: AB2577; MUID:21608550; PMID:11743193
A; Accession: AD3135
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-240 < KUR>
A;Cross-references: UNIPROT:Q8U6V2; UNIPARC:UPI0000164993; GB:AE008689; PIDN:AAL45498.
A; Experimental source: strain C58 (Dupont)
C; Genetics:
A; Gene: huy
A; Map position: linear chromosome
 Query Match
                        39.1%; Score 457.5; DB 2;
                                                   Length 240;
  Best Local Similarity
                        42.1%; Pred. No. 9.1e-29;
 Matches
          98; Conservative
                              41; Mismatches
                                               93;
                                                   Indels
                                                                 Gaps
                                                                        1;
Qу
           1 MRILVINPNSSSALTESVADAAQQVVATGTIISAINPSRGPAVIEGSFDEALATFHLIEE 60
             | | :||||::::|
                              1:1:1
                                        Db
           1 MHIRLINPNSTASMTAQALDSALRVKQKDTHVSAANPVDTPVSIEGQADEAMAVPGLLAE 60
          61 VERAERENPPDAYVIACFGDPGLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVSILPR 120
Qу
                      61 IRKGEGHG-VDAYVIACFDDPGLHAAREVARGPVIGICQAAVQVAMTISRRFSIITTLPR 119
Db
         121 VRKHLHELVRQAGATNRLASIKLPNLGVMAFHEDEHAALETLKQAAKEAVQEDGAESIVL 180
Qу
                                                  - 1 1
                                :: :| |:
                                          | | |
                                              - 1
Db
         120 SIPIIEDLVEDYGAQRYCRKVRAIDLPVLGLEEDPEVAEALLRREIEAAKREDAAEAIIL 179
         181 GCAGMVGFARQLSDELGVPVIDPVEAACRVAESLVALGYQTSKANSYQKPTEK 233
Qу
                      Db
         180 GCAGMSSLCDRLRDATGVPVIDGVTAAIKLAEALVGAGYTTSKVNAYDYPRVK 232
<!--EndFragment-->
```

```
<!--StartFragment-->RESULT 7
S62582
probable hydantoin racemase - fission yeast (Schizosaccharomyces pombe)
C; Species: Schizosaccharomyces pombe
C;Date: 16-May-1996 #sequence_revision 13-Mar-1997 #text change 09-Jul-2004
C; Accession: S62582; T38104
R; Murphy, L.; Niblett, D.; Harris, D.
submitted to the EMBL Data Library, November 1995
A; Reférence number: S62573
A; Accession: S62582
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-238 < MUR>
A;Cross-references: UNIPROT:Q09921; UNIPARC:UPI000013A105; EMBL:Z67998; NID:g1067202;
R; Beck, A.; Reinhardt, R.; Murphy, L.; Niblett, D.; Harris, D.; Barrell, B.G.; Rajandr
submitted to the EMBL Data Library, November 1995
A; Reference number: Z21769
A; Accession: T38104
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: DNA
A; Residues: 1-238 <MU2>
A;Cross-references: UNIPARC:UPI000013A105; EMBL:Z67998; NID:g1067202; PIDN:CAA91957.1;
A; Experimental source: strain 972h-; cosmid c1F7
C; Genetics:
A; Gene: SPAC1F7.10
A; Map position: 1R
                       20.4%; Score 239; DB 2;
 Query Match
 Best Local Similarity 30.1%; Pred. No. 1.4e-11;
          71; Conservative 42; Mismatches 111; Indels
 Matches
                                                            12; Gaps
                                                                        7;
           1 MRILVINPNSSSALTESVADAAQQVVATGTIISAIN-PSRGPAVIEGSFDEALATFHLIE 59
Qу
             : : | || ||
                                                         1 : 1::
           2 IRILVINPNSTVQMTESVKSVLDDCTPPNVQLEYLTCPPEGPKAIECVSDGVRSAAVLMK 61
Db
          60 EVERAERENPP--DAYVIACFGD-PGLDAVKELTDRPVVGVAEAAIHMSSFVAATFSIVS 116
Qу
               Db
          62 YFE----DHPPQVDAFLVSCYSDHPLVTTLRETYRKPCTGIMQASILTALSLGRKVSVVT 117
         117 ILPRVRKHLHELVRQAGATNRL-ASIKLPNLGVMAFHEDEHAALET-LKQAAKEAVQEDG 174
Qу
                    1:: 1::: 1 : :
                                                   1:: 1:1111
Db
         118 TTKRYEPLLTDGIHAMGISDSVFAGIASTGLAPLELDSKPRAEVDALLARTALRAVNEMG 177
         175 AESIVLGCAGMVGFARQLSDELG--VPVIDPVEAACRVAESLVALGYQTSKANSYQ 228
Qу
             : !!! :
                                                          111
         178 ADVICLGCAGMTHMAHVLEKAVGPNIPIIDGTKAGVELLASLVRMNLFTSKQGVYQ 233
<!--EndFragment-->
```